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RADEMAR		1	6 25
	HPS 65 – <u>M.T.</u>	MAKTI	AYDEEARRGL ERGLNALADA
•	HSP 60 – <u>RAT</u>	MLRLPTVLRQ MRPVSRALAP HLTRAYAKDV	KFGADARALM LQGVDLLADA
	HSP 60 - HUMAN	MLRLPTVFRQ MRPVSRVLAP HLTRAYAKDV	
	Consensus	AK	ARGLADA
	Collections		
		26	75
	HPS 65 - M.T.	VKVTLGPKGR NVVLEKKWGA PTITNDGVSI	AKEIELEDPY EKIGAELVKE
:	HSP 60 – <u>RAT</u>	VAVTMGPRGR TVIIEQSWGS PKVTKDGVTV	
·	HSP 60 – <u>HUMAN</u>	VAVTMGPKGR TVIIEQSWGS PKVTKDGVTV	
	HSP 60 - <u>HOMAIN</u>	ANALIMITATION OF THE STATE OF T	
		V-VT-GPKGR -VEWG- PT-DGV	AK-I-L-D-YIGA-LV
•	Consensus		*
		6-7 (31-52 AA)	
		,	
•			
	TIDO CE ACT	76	125
	HPS 65 – <u>M.T.</u>	VAKKTDDVAG DGTTTATVLA QALVREGLRN	
	HSP 60 – <u>RAT</u>	VANNTNEEAG DGTTTATVLA RSIAKEGFER	
•	HSP 60 – <u>HUMAN</u>	VANNTNEEAG DGTTTATVLA RSIAKEGFER	ISKGANPVEI RRGVMLAVDA
			CAND AV
	Consensus	VATAG DGTTTATVLAEG	- 23-GAMPOON - RGAVO-
			21 (121-136 AA)
		126	174
	HPS 65 - <u>M.T.</u>	VTETLIKGAK EVETKEQIAA TAAISA.GDO	SIGDLIAEAM DKVGNEGVIT
	HSP 60 - <i>RAT</i>	VIAELKKOSK PVTTPEEIAQ VATISANGD	K DIGNIISDAM KKVGRKGVIT
	HSP 60 - HUMAN	VIAELKKOSK PVTTPEEIAQ VATISANGDI	A STANTICH IN VICTORIAN
	1101 00		K EIGNIISDAM KKAGKWAII
	Consensus		
	Consensus	VL-KK -V-T-E-IAA-ISA-GD	
	Consensus	VL-KK -V-T-E-IAA-ISA-GD	IGIAM -KVGGVIT
		<u>VL-KK -</u> V-T-E-IAA-ISA-GD	IGIAM -KVGGVIT
,	HPS 65 - <u>M.T.</u>	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTFGLQ LELTEGMRFD KGYISGYFV	IGIAM -KVGGVIT 224 T DPERQEAVLE DPYILLVSK
·	HPS 65 - <u>M.T.</u> HSP 60 - <u>RAT</u>	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTFGLQ LELTEGMRFD RGYISGYFV VKDGKTLNDE LEIIEGMKFD RGYISPYFI	T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK
·	HPS 65 - <u>M.T.</u>	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTFGLQ LELTEGMRFD KGYISGYFV	T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK
·	HPS 65 - <u>M.T.</u> HSP 60 - <u>RAT</u>	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTPGLQ LELTEGMRPD KGYISGYFV VKDGKTLNDE LEIIEGMKPD RGYISPYFI VKDGKTLNDE LEIIEGMKPD RGYISPYFI	T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK
	HPS 65 - <u>M.T.</u> HSP 60 - <u>RAT</u>	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTFGLQ LELTEGMRFD RGYISGYFV VKDGKTLNDE LEIIEGMKFD RGYISPYFI	T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK
	HPS 65 - <u>M.T.</u> HSP 60 - <u>RAT</u> HSP 60 - <u>HUMAN</u>	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTPGLQ LELTEGMRPD KGYISGYFV VKDGKTLNDE LEIIEGMKPD RGYISPYFI VKDGKTLNDE LEIIEGMKPD RGYISPYFI VT LEEGM-PD -GYIS-YF-	224 T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK N TSKGQKCEFQ DAYVLLSEKK
•	HPS 65 - <u>M.T.</u> HSP 60 - <u>RAT</u> HSP 60 - <u>HUMAN</u>	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTPGLQ LELTEGMRPD KGYISGYFV VKDGKTLNDE LEIIEGMKPD RGYISPYFI VKDGKTLNDE LEIIEGMKPD RGYISPYFI	T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK
•	HPS 65 - <u>M.T.</u> HSP 60 - <u>RAT</u> HSP 60 - <u>HUMAN</u>	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTFGLQ LELTEGMRFD KGYISGYFV VKDGKTLNDE LEIIEGMRFD RGYISPYFI VKDGKTLNDE LEIIEGMRFD RGYISPYFI VT LEEGM-FD -GYIS-YF- 31 (181-196 AA)	224 T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK N TSKGQKCEFQ DAYVLLSEKK
	HPS 65 – <u>M.T.</u> HSP 60 – <u>RAT</u> HSP 60 – <u>HUMAN</u> Consensus	VL-KK -V-T-E-IAA-ISA-GD- 175 VEESNTFGLQ LELTEGMRFD RGYISGYFV VKDGKTLNDE LEIIEGMKFD RGYISPYFI VKDGKTLNDE LEIIEGMKFD RGYISPYFI VT LEEGM-FD -GYIS-YF- 31 (181-196 AA)	224 T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK N TSKGQKCEFQ DAYVLLSEKKQ D-Y-LLK 36 (211-226 AA)
	HPS 65 – <u>M.T.</u> HSP 60 – <u>RAT</u> HSP 60 – <u>HUMAN</u> Consensus	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTFGLQ LELTEGMRFD KGYISGYFV VKDGKTLNDE LEIIEGMRFD RGYISPYFI VKDGKTLNDE LEIIEGMRFD RGYISPYFI VT LEEGM-FD -GYIS-YF- 31 (181-196 AA) 225 VSTVKDLLPL LEKVIGAGKP LLIIAEDVE	224 T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK N TSKGQKCEFQ DAYVLLSEKKQ D-Y-LLK 36 (211-226 AA) 274 G EALSTLVVNK IRGTYKSVAV
	HPS 65 - <u>M.T.</u> HSP 60 - <u>RAT</u> HSP 60 - <u>HUMAN</u> Consensus HPS 65 - <u>M.T.</u> HSP 60 - <u>RAT</u>	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTPGLQ LELTEGMRFD RGYISGYFV VKDGKTLNDE LEIIEGMKFD RGYISPYFI VKDGKTLNDE LEIIEGMKFD RGYISPYFI VT LEEGM-FD -GYIS-YF- 31 (181-196 AA) 225 VSTVKDLLPL LEKVIGAGKP LLIIAEDVE ISSVQSIVPA LEIANAHRKP LVIIAEDVE	224 T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK N TSKGQKCEFQ DAYVLLSEKKQ D-Y-LLK 36 (211-226 AA) 274 G EALSTLVVNK IRGTYKSVAV LKVGLQVVAV
	HPS 65 – <u>M.T.</u> HSP 60 – <u>RAT</u> HSP 60 – <u>HUMAN</u> Consensus	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTFGLQ LELTEGMRFD KGYISGYFV VKDGKTLNDE LEIIEGMRFD RGYISPYFI VKDGKTLNDE LEIIEGMRFD RGYISPYFI VT LEEGM-FD -GYIS-YF- 31 (181-196 AA) 225 VSTVKDLLPL LEKVIGAGKP LLIIAEDVE	224 T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK N TSKGQKCEFQ DAYVLLSEKKQ D-Y-LLK 36 (211-226 AA) 274 G EALSTLVVNK IRGTYKSVAV LKVGLQVVAV
	HPS 65 - <u>M.T.</u> HSP 60 - <u>RAT</u> HSP 60 - <u>HUMAN</u> Consensus HPS 65 - <u>M.T.</u> HSP 60 - <u>RAT</u> HSP 60 - <u>RAT</u>	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTPGLQ LELTEGMRFD KGYISGYFV VKDGKTLNDE LEIIEGMKFD RGYISPYFI VKDGKTLNDE LEIIEGMKFD RGYISPYFI VT LEEGM-FD -GYIS-YF- 31 (181-196 AA) 225 VSTVKDLLPL LEKVIGAGKP LLIIAEDVE ISSVQSIVPA LEIANAHRKP LVIIAEDVE ISSIQSIVPA LEIANAHRKP LVIIAEDVE	224 T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK N TSKGQKCEFQ DAYVLLSEKKQ D-Y-LLK 36 (211-226 AA) 274 G EALSTLVUNK IRGTYKSVAV G EALSTLVLNR LKVGLQVVAV LKVGLQVVAV
	HPS 65 - <u>M.T.</u> HSP 60 - <u>RAT</u> HSP 60 - <u>HUMAN</u> Consensus HPS 65 - <u>M.T.</u> HSP 60 - <u>RAT</u>	VL-KK -V-T-E-IAA-ISA-GD 175 VEESNTPGLQ LELTEGMRFD RGYISGYFV VKDGKTLNDE LEIIEGMKFD RGYISPYFI VKDGKTLNDE LEIIEGMKFD RGYISPYFI VT LEEGM-FD -GYIS-YF- 31 (181-196 AA) 225 VSTVKDLLPL LEKVIGAGKP LLIIAEDVE ISSVQSIVPA LEIANAHRKP LVIIAEDVE	224 T DPERQEAVLE DPYILLVSSK N TSKGQKCEFQ DAYVLLSEKK N TSKGQKCEFQ DAYVLLSEKK N TSKGQKCEFQ DAYVLLSEKK Q D-Y-LLK 36 (211-226 AA) 274 G EALSTLVUNK IRGTYKSVAV G EALSTLVLNR LKVGLQVVAV LKVGLQVVAV

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FIG. 1A

40 (236-251 AA)

45 (265-280 AA)

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HPS 65 – <u>M.T.</u> HSP 60 – <u>RAT</u> HSP 60 – <u>HUMAN</u>	KAPGPGDNRK NQLKDMAIAT	323 G GGQVISEE.V GLTLENADLS LLGKARKVVV G GGAVFGEEGL NLNLEDVQAH DLGKVGEVIV G GGAVFGEEGL TLNLEDVQPH DLGKVGEVIV
Consensus	KAPGFGD-RKL-DMAI-1	GG-VEEL-LELGKV-V
HPS 65 – <u>M.T.</u> HSP 60 – <u>RAT</u> HSP 60 – <u>HUMAN</u> Consensus	TKDDAMLLKG KGDKAHIEKF TKDDAMLLKG KGDKAQIEKF	373 R VAQIRQEIEN SDSDYDREKL QERLAKLAGG R IQEITEQLDI TTSEYEKEKL NERLAKLSDG R IQEIIEQLDV TTSEYEKEKL NERLAKLSDG
Consensus	TKD=====G =GD===I==F	I
	•	59 (349-364 AA)
HPS 65 – <u>M.T.</u> HSP 60 – <u>RAT</u> HSP 60 – <u>HUMAN</u> Consensus	VAVLKVGGTS DVEVNEKKDE	423 R IEDAVRNAKA AVEEGIVAGG GVTLLQAAPT R VTDALNATRA AVEEGIVLGG GCALLRCIPA R VTDALNATRA AVEEGIVLGG GCALLRCIPA RDAA AVEEGIV-GG GLLP-
HPS 65 – <u>M.T.</u> HSP 60 – <u>RAT</u> HSP 60 – <u>HUMAN</u> Consensus	LDSLXPANED QKIGIEIIKF LDSLTPANED QKIGIEIIKF	472 7 ALEAPLKQIA FNSGLEPGVV AEKVRNLPAG R ALKIPAMTIA KNAGVEGSLI VEKILQSSSE R TLKIPAMTIA KNAGVEGSLI VEKIMQSSSELPIA -N-G-EEK
HPS 65 – <u>M.T.</u> HSP 60 – <u>RAT</u> HSP 60 – <u>HUMAN</u> Consensus	VGYDAMLGDF VNMVEKGIII VGYDAMAGDF VNMVEKGIII	522 PVKVTRSALQ NAASIAGLFL TTEAVVADKP PTKVVRTALL DAAGVAPLLT TAEAVVTEIP PTKVVRTALL DAAGVASLLT TAEVVVTEIP P-KV-R-ALAAA-L T-E-VVP
		84 (499-514 AA)
HPS 65 – <u>M.T.</u> HSP 60 – <u>RAT</u> HSP 60 – <u>HUMAN</u>	523 540 EKEKASVPGG GDMGGMDF KEEKDPGM GAMGGMGGGM KEEKDPGM GAMGGMGGGM	GGGMF FIG. 1B

--EK---PG- G-MGGM----

Consensus

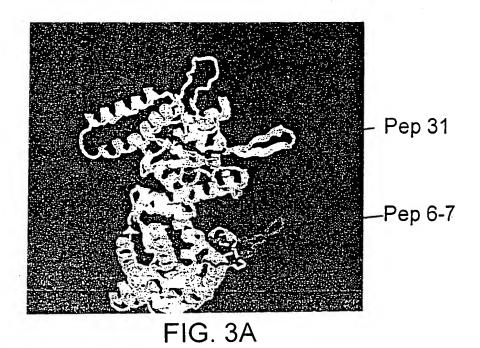
Appln Nr 9/847,637 Page ? 7
Applicant, J: Yaakov Naparstek et al.
NOVEL AMINO ACID SEQUENCES, DNA ENCODING THE
AMINO ACID SEQUENCES, ANTIBODIES DIRECTED
AGAINST SUCH SEQUENCES AND THE DIFFERENT USES
THEREOF

peptide 31 (181-197 AA)

peptides 6-7 (31-52 AA)

FIG. 2

THEREOF



Pep 31 Pep 6-7

FIG. 3B



Appln N⁻ 9/847.637 Page 5 7 Applican_{N-2}: Yaakov Naparstek et al. NOVEL AMINO ACID SEQUENCES, DNA ENCODING THE AMINO ACID SEQUENCES, ANTIBODIES DIRECTED AGAINST SUCH SEQUENCES AND THE DIFFERENT USES THEREOF

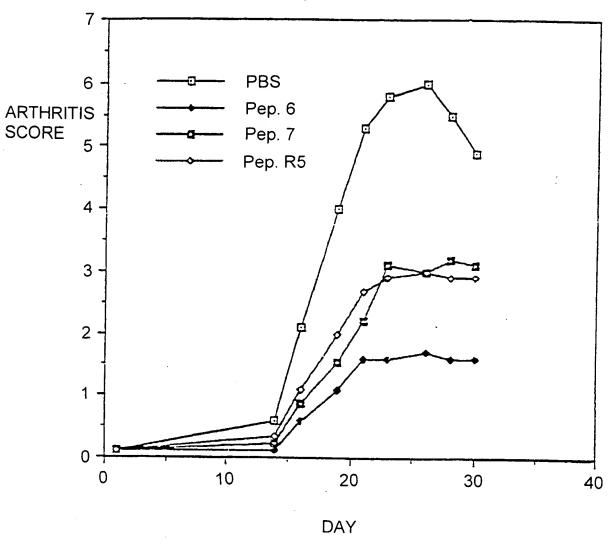


FIG. 4

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NOVEL AMINO ACID SEQUENCES, DNA ENCODING THE AMINO ACID SEQUENCES, ANTIBODIES DIRECTED AGAINST SUCH SEQUENCES AND THE DIFFERENT USES

THEREOF



The "Protective" Motif

ر ال \geq × ¥ Ш >1 Z \simeq G ¥ Д G HSP Peptide 6- (31-46)

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HSP Peptide 5- (36-55)

Rat

Common Motif

HSP Peptide 7- (37-52)

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FIG. 5

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NOVEL AMINO ACID SEQUENCES, DNA ENCODING The

AMINO ACID SEQUENCES, ANTIBODIES DIRECTED

AGAINST SUCH SEQUENCES AND THE DIFFERENT USES **THEREOF**

